RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information

Center, (STIC) detected errors when processing the following computer read tole

form:

Application Serial Number: 09/403, 86/

Source: 35.35.55 - 30.55 1653

Date Processed by STIC: 9-19-00

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR FURTHER INFORMATION, PLEASE TELEPHONE MARK SPENGER.
703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/403,861

DATE: 09/19/2000 TIME: 18:40:44

RECEIVED SEP 29 2000

TECH CENTER 1600/280

Does Not Comply

Corrected Diskette Needed

Input Set : A:\RICCARD1.txt

Output Set: N:\CRF3\09192000\I403861.raw

```
SEQUENCE LISTING
4 (1) GENERAL INFORMATION:
        (i) APPLICANT: RICCARDI, Carlo
       (ii) TITLE OF INVENTION: INTRACELLULAR MODULATORS OF APOPTIC CELL
                               DEATH PATHWAYS
      (iii) NUMBER OF SEQUENCES: 15
      (iv) CORRESPONDENCE ADDRESS:
13
14
15
```

(A) ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C. (B) STREET: 624 Ninth Street, N.W., Suite 300 (C) CITY: Washington

(D) STATE: D.C. (E) COUNTRY: USA (F) ZIP: 20001

21 (V) COMPUTER READABLE FORM: (A) MEDIUM TYPE: Floppy disk (B) COMPUTER: IBM PC compatible

(C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (vi) CURRENT APPLICATION DATA:

27 (A) APPLICATION NUMBER: US/09/403,861 C--> 28

C--> 29 (B) FILING DATE: 11-Feb-2000 C--> 35 (vii) PRIOR APPLICATION DATA:

32 (A) APPLICATION NUMBER: PCT/EP98/02490

33 (B) FILING DATE: 27-APR-1998

(A) APPLICATION NUMBER: EP 97107033.9 36 (B) FILING DATE: 28-APR-1997 37

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: YUN, Allen C. 40

(B) REGISTRATION NUMBER: 37,971 41 42 (C) REFERENCE/DOCKET NUMBER: RICCARDI=1

44 (ix) TELECOMMUNICATION INFORMATION: 45 (A) TELEPHONE: 202-628-5197

(B) TELEFAX: 202-737-3528

ERRORED SEQUENCES

17 18

19

24 25

39

```
544 (2) INFORMATION FOR SEQ ID NO: 14:
                                             -> See next page
546
         (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 26 amino acids — (B) TYPE: amino acid
547
548
549
              (D) TOPOLOGY: linear
551
        (ii) MOLECULE TYPE: peptide
        (ix) FEATURE:
             (D) OTHER INFORMATION:/note= "the N-terminus is modified by an
555 acetyl group; the C-terminus is modified with a-(4-methyl-coumaryl-
556 7-amide)"
```

DATE: 09/19/2000 TIME: 18:40:44

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/403,861

Input Set : A:\RICCARD1.txt
Output Set: N:\CRF3\09192000\1403861.raw

	• • • • • • • • • • • • • • • • • • • •	
563 565 566 567 568 570 572 573	Asp Glu Val Asp (2) INFORMATION FOR SEQ ID NO: 15: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear	
576	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:	200(0) (2/0 (0:5/2):2
	Tyr Val Ala Asp	
E7 378	Tyr val Ala Asp	26 is listed as
		length, only 4
8		amino acids are shown.

VERIFICATION SUMMARY DATE: 09/19/2000 PATENT APPLICATION: US/09/403,861 TIME: 18:40:45

Input Set : A:\RICCARD1.txt

Output Set: N:\CRF3\09192000\I403861.raw

L:28 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:29 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:35 M:220 C: Keyword misspelled or invalid format, [(vii) PRIOR APPLICATION DATA:]
L:270 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:5
L:394 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=7, Value=[pepetide]
L:560 M:203 E: No. of Seq. differs, LENGTH:Input:26 Found:4 SEQ:14
L:578 M:203 E: No. of Seq. differs, LENGTH:Input:26 Found:4 SEQ:15